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09/470,276 12/22/99 KOLODNER R 157/47483-C

NIXON PEABODY LLP
101 FEDERAL ST
BOSTON MA 02115

HM12/0206

EXAMINER

FREDMAN, J

ART UNIT

PAPER NUMBER

1655

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Please find below and/or attached an Office communication concerning this application or proceeding.

Commissioner of Patents and Trademarks

Office Action SummaryApplication No.
09/470,276

Applicant(s)

Kolodner et al

Examiner

Jeffrey Fredman

Group Art Unit

1655☒ Responsive to communication(s) filed on Jan 2, 2001☐ This action is **FINAL**.☐ Since this application is in condition for allowance except for formal matters, prosecution as to the merits is closed in accordance with the practice under *Ex parte Quayle*, 1935 C.D. 11; 453 O.G. 213.

A shortened statutory period for response to this action is set to expire three month(s), or thirty days, whichever is longer, from the mailing date of this communication. Failure to respond within the period for response will cause the application to become abandoned. (35 U.S.C. § 133). Extensions of time may be obtained under the provisions of 37 CFR 1.136(a).

Disposition of Claims☒ Claim(s) 1-38 is/are pending in the application.Of the above, claim(s) 1 and 13-38 is/are withdrawn from consideration.☐ Claim(s) _____ is/are allowed.☒ Claim(s) 2-12 is/are rejected.☐ Claim(s) _____ is/are objected to.☐ Claims _____ are subject to restriction or election requirement.**Application Papers**☐ See the attached Notice of Draftsperson's Patent Drawing Review, PTO-948.☐ The drawing(s) filed on _____ is/are objected to by the Examiner.☐ The proposed drawing correction, filed on _____ is ☐ approved ☐ disapproved.☐ The specification is objected to by the Examiner.☐ The oath or declaration is objected to by the Examiner.**Priority under 35 U.S.C. § 119**☐ Acknowledgement is made of a claim for foreign priority under 35 U.S.C. § 119(a)-(d).☐ All ☐ Some* ☐ None of the CERTIFIED copies of the priority documents have been☐ received.☐ received in Application No. (Series Code/Serial Number) _____.☐ received in this national stage application from the International Bureau (PCT Rule 17.2(a)).

*Certified copies not received: _____

☒ Acknowledgement is made of a claim for domestic priority under 35 U.S.C. § 119(e).**Attachment(s)**☒ Notice of References Cited, PTO-892☒ Information Disclosure Statement(s), PTO-1449, Paper No(s). 5☐ Interview Summary, PTO-413☐ Notice of Draftsperson's Patent Drawing Review, PTO-948☐ Notice of Informal Patent Application, PTO-152

--- SEE OFFICE ACTION ON THE FOLLOWING PAGES ---

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DETAILED ACTION

Election/Restriction

1. Applicant's election with traverse of Group II, claims 2-12 and SEQ ID Nos: 1, 3-6, 27-30 and 53, in Paper No. 9 is acknowledged. The traversal is on the ground(s) that each of the remaining sequences are human sequences. This is not found persuasive because this is not a reason to overcome a restriction requirement. Presumably the intention is to argue that the sequences lack patentable distinctness, however, the fact that sequences from two different vertebrate species are included is evidence that the sequences are distinct. With regard to burden, it would represent a significant burden to search each additional sequence.

The requirement is still deemed proper and is therefore made FINAL.

General

2. Claims 2-12 utilize the transitional term "having". Because this term lacks any particular meaning in the patent literature, the examiner will interpret "having" as being equivalent in scope to the open term "comprising".

Claim Rejections - 35 USC § 112

3. The following is a quotation of the first paragraph of 35 U.S.C. 112:

The specification shall contain a written description of the invention, and of the manner and process of making and using it, in such full, clear, concise, and exact terms as to enable any person skilled in the art to which it pertains, or with which it is most nearly connected, to make and use the same and shall set forth the best mode contemplated by the inventor of carrying out his invention.

4. Claims 5-11 are rejected under 35 U.S.C. 112, first paragraph, as containing subject matter which was not described in the specification in such a way as to reasonably convey to one

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skilled in the relevant art that the inventor(s), at the time the application was filed, had possession of the claimed invention.

The current claims are drawn to one of two broad genus, claims 5-9 being drawn to a genus comprising any "unique fragment" of SEQ ID NO: 1 and claim 10-11 being drawn to any primers which permit synthesis of a human mismatch repair gene, particularly hMSH5. This large genus is represented in the specification by only the named SEQ ID Nos. Thus, applicant has express possession of only one full length nucleic acid species and shows no fragments which are demonstrably unique and shows multiple primers in a genus which comprises hundreds of millions of different possibilities. The written description guidelines note regarding such genus/species situations that "Satisfactory disclosure of a ``representative number" depends on whether one of skill in the art would recognize that the applicant was in possession of the necessary common attributes or features of the elements possessed by the members of the genus in view of the species disclosed." (See: Federal Register: December 21, 1999 (Volume 64, Number 244), revised guidelines for written description.) Here, no common element or attributes of the sequences are disclosed, not even the presence of certain domains. No structural limitations or requirements which provide guidance on the identification of sequences which meet these functional limitations is provided.

Further, these claims encompass alternately spliced versions of the proteins, allelic variants including insertions and mutations, inactive precursor proteins which have a removable amino terminal end, and only specific nucleic acid sequences have been provided. No written description

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of alleles, of upstream or downstream regions containing additional sequence, or of alternative splice variants has been provided in the specification.

It is noted that in Fiers v. Sugano (25 USPQ2d, 1601), the Fed. Cir. concluded that

"...if inventor is unable to envision detailed chemical structure of DNA sequence coding for specific protein, as well as method of obtaining it, then conception is not achieved until reduction to practice has occurred, that is, until after gene has been isolated...conception of any chemical substance, requires definition of that substance other than by its functional utility."

In the instant application, only the nucleic acid and inherent amino acid sequence of the disclosed SEQ ID Nos are described. Also, in Vas-Cath Inc. v. Mahurkar (19 USPQ2d 1111, CAFC 1991), it was concluded that:

"...applicant must also convey, with reasonable clarity to those skilled in art, that applicant, as of filing date sought, was in possession of invention, with invention being, for purposes of "written description" inquiry, whatever is presently claimed."

In the application at the time of filing, there is no record or description which would demonstrate conception or written description of any nucleic acids acids which comprise "unique fragments" or which are modified by addition, insertion, deletion, substitution or inversion with the disclosed SEQ ID Nos.

Claim Rejections - 35 USC § 102

5. The following is a quotation of the appropriate paragraphs of 35 U.S.C. 102 that form the basis for the rejections under this section made in this Office action:

A person shall be entitled to a patent unless --

(b) the invention was patented or described in a printed publication in this or a foreign country or in public use or on sale in this country, more than one year prior to the date of application for patent in the United States.

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6. Claims 2-6, 8 and 9 are rejected under 35 U.S.C. 102(b) as being anticipated by Sargent et al (EMBO J. (1989) 8(8):2305-2312).

Sargent teaches cosmid vectors which are transformed into *E. Coli* host cells (page 2311, column 2) which cosmid vectors comprise a double stranded nucleic acid (necessarily including sense and antisense strands) that includes the "G7" gene (page 2306, figure 1). Within figure 1, several cosmid vectors overlap the "G7" gene including F9N, F12M, FMC, FMEa and EL3. The "G7" gene is inherently found to be the MSH5 gene claimed. With regard to points of similarity, the "G7" gene is located on chromosome 6p21.3 as is the MSH5 gene, and the following is a partial sequence comparison of MSH5 and "G7", where MSH5 is the Qy sequence and "G7" is the database sequence.

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Query Match          89.5%; Score 2596; DB 68; Length 3998;
Best Local Similarity 97.1%; Pred. No. 0;
Matches 2675; Conservative 0; Mismatches 30; Indels 51; Gaps 1;

Qy 184  tgcggccacagggcgttcagacccctctttccaaaggagcctccagctcatgtggcctcc 243
    ||| | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
Db 6  TCGGTTCAGCGGGGCGTTCTCCACGTGTAGCGACTCAGAGCTCCAGCTCATGGCTCC 65

Qy 244  ttaggagcgaccccaaggaggacaccgcaggagaccgagacctggggcggtccctccgtt 303
    ||||| | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
Db 66  TTAGGAGCGAACCACAGGAGGACACCGCAGGGACCGAGACCTGGGGCGGCTCCTCCGGC 125

Qy 304  ttccccagccgggccccagtgccggggccccaggaggcccgaggaggaggaaagtcgaggag 363
    ||||| | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
Db 126  TTCCCCAGCCCGGCCCGAGTGCCGGGCCCCAGGGAGGCCGAGGAGGAGGAAGTCGAGGAG 185

Qy 364  gaggaggagctggccgagatccactctgtgtgtgtgtggaattcaggatacttgggcatt 423
    ||||| | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
Db 186  GAGGAGGAGCTGGCCGAGATCCATCTGTGTGTGCTGTGGAATTCAGGATACTTGGGCTT 245

Qy 424  gcctactatgatactagtgaactccactatccacttcgatgocagatgccccagacacagag 483
    ||||| | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
Db 246  GCCTACTATGATACTAGTGACTCCACTATCCACTTCATGCCAGATGCCCCAGACCCAGAG 305

Qy 484  agcctcaagcttctccagagagttctggatgagatcaatcccagctctgttggtaacaggt 543
    ||||| | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
Db 306  AGCCTCAAGCTTCTCCAGAGAGTTCTGGATGAGATCAATCCCAGTCTGTTGTTAAGGT 365

Qy 544  gccaaacaggatgagaatatgactegattctctgggaaagcttgccctccagagacacaga 603
    ||||| | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
Db 366  GCCAAACAGGATGAGATATGACTCGATTCTCTGGGAAGCTTCGCTCCAGGAGCAGGA 425

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Qy 604 gagcctaaaagacctgaatcatatcttttggcaagtggtgattttggtctggagataaagc 663
Db 426 GAGCCTAAAAGACCTGAAATCATATTTTGGCCAAAGTGTGGATTTTGGTCTGGAGATAAGC 485

Qy 664 aaacaacgctccttttctggaactactccttctccccaagaagccactgactgccactgag 723
Db 486 AAACAACGCTCCTTTCTGGAACCTACTCCTTTCATCCACAGACGCCATGACTGCCACTGAG 545

Qy 724 aaaatcctcttctcctctcttccattattccctttgactgctcctcaca----- 771
Db 546 AAAATCCTCTTCTCTCTCTCCATTATTCCTTTGACTGCCTCCTCACACCCOCCAGGAGAT 605

Qy 772 -----gttcgagcaattggagggtg 792
Db 606 TTAAGATTACCCCGATTCCGACTGCTGATCCOCTCCAGGTTCCGAGCACTTGGAGGGCTG 665

Qy 793 ctgaagttcctgggtggaagaagaatcggggtgaactggaagactataatgtcagcgctc 852
Db 666 CTGAAGTTCTCGGTCGAAGAAGAAATCGGGGTGAAGTGGAAAGACTATAATGTGACGCTC 725

Qy 853 cccatcctgggctttaagaatttatgttgactcatctggtgaacatagatcaagacact 912
Db 726 CCCATCTCGGGCTTTAAGAAATTTATGTTGACTCATCTGGTGAACATAGATCAAGACACT 785

Qy 912 tacagtgttctacagatttttaagagtgagttcaccocctcaagtgtacaaagtgccagct 972
Db 786 TACAGTGTCTACAGATTTTAAAGAGTGAGTCTCACOCTCAGTGTACAAAGTGGCCAGT 845

Qy 973 ggactgaaggagggtcagcctcttttggaatcctcaacagatgccactgtgaagtgggga 1032
Db 846 GGAAGTGAAGGAGGGGCTCAGCCTCTTTGGAATCCTCAACAGATGCCACTGTAAAGTGGGA 905

Qy 1033 gagaagctgctcaggctatggttcacacgctgcagactcatgaactgggggagctcagttct 1092
Db 906 GAGAAGCTGCTCAGGCTATGGTTCAAGCTCCGACTCATGAOCTGGGGAGCTCAGTTCT 965

Qy 1093 cgtctggagctcattcagttttctctgctgcccaagaatctggacatggctcagatgctg 1152
Db 966 CGCTGGAGCTCATTGAGTTTCTCTGCTGCCOCCAGAATCTGGACATGGCTCAGATGCTG 1025

Qy 1153 catcgctcctgggtcacatcaagaagctgcttgattctgaaagcattgaagttgtcc 1212
Db 1026 CATCGCTCCTGGGTCAATCAAGAACGTTGCTCTGATTCTGAAACGATGAAGTTGTCTC 1085

Qy 1213 caccacaaggtcagcagctggcaggttctctacaagactgtgtacagtgccctgggctg 1272
Db 1086 CACRCCAAAGTCAAGCACTGGCAGGTTCTCTCAAGACTGTGTACAGTGCCCTGGGCTG 1145

Qy 1273 agggatgctcgcgctcctgcgcagctccatccagctctttogggacattgcccaagag 1332
Db 1146 AGGGATGCTGCGCTCCTGCGCGAGTCCATCCAGCTCTTTGGGACATTGCCCAAGAG 1205

Qy 1333 tctctgatgactgcacatatcgccagctcattgggaaagttagtgactttgagggc 1392
Db 1206 TTCTGATGACCTGCACATATCGCGAGCCTCATGGGAAAGTAGTGACTTTGAGGGC 1265

Qy 1393 agccttgctgaaatcgcttcacagctcctcccaacatagatcctgaaattgatgagaaa 1452
Db 1266 AGCCTTGCTGAAATTCGCTTCAGAGCTCCTCCCAACATAGATCCTGAAATTGATGAGAAA 1325

Qy 1453 aagcgaagactgattggagcttcccaagtttcttaactgaggttgcccgcaaggagctggag 1512
Db 1326 AAGCGAAGACTGATGGGACTTCCCAAGTTTCCTTACTGAGGTTGCCCGCAAGAGCTGGAG 1385

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Qy 1513 aatctggaactcccgattatcctcatgcaagtgaatcatcatcctctgattggcttcctt 1572
Db 1386 AATCTGGACTCCCGATTTCCTTCATGCAAGTGCATCTACATCOCTCTGATTGGCTTCCTT 1445

Qy 1573 atttctattcccgctgcttccatggttagagggcagtgactttgagattaatggactg 1632
Db 1446 CTTTCTATTCCCGCGCTGCTTCCATGSGTAGAGGCGAGTGACTTTGAGATTAAATGGACTG 1505

Qy 1633 gacttcactgtttctctcagaggagaagctgcactatcgtagtgcccgaaacaaaggagctg 1692
Db 1506 GACTTCATGTTTCTCTCAGAGGAGAAGCTGCACATCTAGTGCCCGAACCAAGGAGCTG 1565

Qy 1693 gatgcattgtctgggggacctgcactgcagatccgggaccaggagacgtgctgatgtac 1752
Db 1566 GATGCATTGTCTGGGGACCTGCACCTGCGAGATCCGGGACCAGGAGACGCTGCTGATGTAC 1625

Qy 1753 cagctacagtgccaggtgtctggcagagcagctgtcttaaccgcagattatggaaacttgcc 1812
Db 1626 CAGCTACAGTGCCAGGTGTGGCAGAGCAGCTGTCTTAACCCGAGTATTGGACCTTGCC 1685

Qy 1813 tcccgcttggaagctcctgctgctctctgocagtgctgcccgggactatggctactcaagg 1872
Db 1686 TCCCGCTTGGAAGCTCCTGCTGGCTCTTGCACATGCTGCCCGGACATGGCTACTCAAG 1745

Qy 1873 ccgggttactcccaacaagtctctgggtgaagaatccagaatggagacatcctctgatg 1932
Db 1746 CCGCTTACTCCCAACAAGTCTCTGGGGTACGAATCCAGAATGGCAGACATCCTCTGATG 1805

Qy 1933 gaactctgtgcccgaacctttgtgcccaactccacagaatgtgggtggggacaaaggagg 1992
Db 1806 GAACCTCTGTGCCGAACCTTTGTGCCCAACTCCACAGAATGTGGTGGGACAAAGGAGG 1865

Qy 1993 gtcaaaagtcacactggaacccaactcatcagggaagagcatataacctcaaacaggtaggc 2052
Db 1866 GTCAAAAGTCATCACTGGAACCAACTCATCAGGGAAGAGCATATACCTCAAAACAGTAGGC 1925

Qy 2053 ttgatcacattcatgcccctggtaggcagctttgtgccagcagaggaggccgaattggg 2112
Db 1926 TTGATCACATTTCATGGCCCTGGTAGGCAGCTTTGTGCCAGCAGAGGAGCCGAATTTGGG 1985

Qy 2113 gcaagttagcagccatcttcacacgaattcatagctgggaatccatctccttggcctctcc 2172
Db 1986 GCAGTAGACGCCATCTTCACAGGAATTCATAGCTGCGAATCCATCTCCCTTGGCCTCTCC 2045

Qy 2173 accttcattgatgcacctcaaccagggtggcgaaagcagtgaaacaatgccactgcacatgct 2232
Db 2046 ACCTTCATGATCGAAGCTCAACCAAGGTGGCGAAAGCAGTGAACAATGCCACTGCACATGCTG 2105

Qy 2233 ctggtccttattgatgaatttggaaagggaacccaacacaggtggatgggctcggcctctctg 2292
Db 2106 CTGTCCTCTATTGATGAATTGGAAAGGGAACCAACACGGTGGATGGGCTCGGCTCTCTG 2165

Qy 2293 ggcgtgtgtctccgacactggctggcagctgggaacccacatgcccacacatcctttgigggcc 2352
Db 2166 GCGCTGTGCTCCGACACTGCTGCTGCAAGTGGACCCACATGCCCCACAGTCTTTGTGGCC 2225

Qy 2353 accaaacttctgagccttgttcagctacaactgctgcccacaaaggcccccctgggtgcagtat 2412
Db 2226 ACCAACTTCTTGAGCCTTGTTTCAGCTACAACCTGCTGCCACAGGGGCCCTGGTGAGATG 2285

Qy 2413 ttgacatggagacactgtgaggatggcaacagatcttctcttctatcaggtttggcaa 2472
Db 2286 TTGACCATGGAGACCTGTGAGGATGGCAACGATCTTGTCTCTCTATCAGGTTTGGGAA 2345

Qy 2473 ggtgttgogaaggccagccatgcctcccaacacagctgcccaggtgggcttctgacaaag 2532

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Db 2346 GGTGTTGCGAAGGCCAGCCATGCTCCACACAGCTGCCAGGCTGGGCTCCTGACAAG 2405
Qy 2533 ctgtggctogtggcaaggaggtctcagatttgatcgcagtggaagaaacccatcaagcct 2592
Db 2406 CTTGTGGCTCSTGGCAAGSAGGTCTCAGACTTGATCCGCAGTGGAAACCCATCAAGCCT 2465
Qy 2593 gtcaaggatttgctaaagaagaacaaatggaaattgcccagacattagtggataagttt 2652
Db 2466 GTCAGGATTTGCTAAAGAAGAAOCAAATGGAAAATYGCCAGACATTAGTGGATAAGTTT 2525
Qy 2653 atgaaactggatttggaagatcctaaccctggacttgaacgttttcatgagccaggaagt 2712
Db 2526 ATGAAACTGGATTGGAGATCTTAACCTGGACTTGAACGTTTTCATGAGCCAGGAAGT 2585
Qy 2713 ctgcctgtgcaccagacatcctctgagagtccttcagagtgctctcccagcctcctcag 2772
Db 2586 TGCTGCTGCTGCTACAGCATCTCTGAGATCCTTCCAGTGTCTCCCCAGCCTCCTGAG 2645
Qy 2773 actcoggtgggctgcatgcctctttgtttccttatctccctcagacgcagagttttta 2832
Db 2646 ACTCCGGTGGGCTGCCATGCCCTCTTTGTTTCTTATCTCCTCAGAGCAGAGTTTITA 2705
Qy 2833 gtttctctagaaattttgtttcattattaggaataaagtgtttatttgaagaaaaaa 2888
Db 2706 GTTCTCTAGAAATTTTGTTCATATTAGGAATAAGTTTATTTTGAAGAAAGATA 2761

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Therefore, it clearly appears that the Sargent cosmid vectors anticipate the current claims as inherently comprising the MSH5 gene sequence.

Claim Rejections - 35 USC § 103

7. The following is a quotation of 35 U.S.C. 103(a) which forms the basis for all obviousness rejections set forth in this Office action:

(a) A patent may not be obtained though the invention is not identically disclosed or described as set forth in section 102 of this title, if the differences between the subject matter sought to be patented and the prior art are such that the subject matter as a whole would have been obvious at the time the invention was made to a person having ordinary skill in the art to which said subject matter pertains. Patentability shall not be negated by the manner in which the invention was made.

This application currently names joint inventors. In considering patentability of the claims under 35 U.S.C. 103(a), the examiner presumes that the subject matter of the various claims was commonly owned at the time any inventions covered therein were made absent any evidence to the contrary. Applicant is advised of the obligation under 37 CFR 1.56 to point out the inventor

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and invention dates of each claim that was not commonly owned at the time a later invention was made in order for the examiner to consider the applicability of 35 U.S.C. 103(c) and potential 35 U.S.C. 102(f) or (g) prior art under 35 U.S.C. 103(a).

8. Claims 10-12 are rejected under 35 U.S.C. 103(a) as being unpatentable over Albertella (Genomics (1996) 36:240-251) in view of Stratagene Catalog (1988) p. 39.

Albertella teaches a number of primers which function to amplify the "G7" gene region (page 241, column 2, subheading "reverse-transcription-PCR" to page 242, column 1). As discussed above, "G7" is inherently found to be the MSH5 gene.

Albertella does not teach placement of these reagents into a kit format, nor the specific SEQ ID Nos: 3-50.

Stratagene catalog teaches a motivation to combine reagents into kit format (page 39).

It would have been *prima facie* obvious to one having ordinary skill in the art at the time the invention was made to combine the primers of Albertella into a kit format as discussed by Stratagene catalog since the Stratagene catalog teaches a motivation for combining reagents of use in an assay into a kit, "Each kit provides two services: 1) a variety of different reagents have been assembled and pre-mixed specifically for a defined set of experiments. Thus one need not purchase gram quantities of 10 different reagents, each of which is needed in only microgram amounts, when beginning a series of experiments. When one considers all of the unused chemicals that typically accumulate in weighing rooms, desiccators, and freezers, one quickly realizes that it is actually far more expensive for a small number of users to prepare most buffer

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solutions from the basic reagents. Stratagene provides only the quantities you will actually need, premixed and tested. In actuality, the kit format saves money and resources for everyone by dramatically reducing waste. 2) The other service provided in a kit is quality control" (page 39, column 1).

It would have been *prima facie* obvious to one having ordinary skill in the art at the time the invention was made to identify functionally equivalent primers and probes selected from the sequences disclosed by Albertella for detection of the "G7" gene.

In the recent court decision *In Re Deuel* 34 USPQ 2d 1210 (Fed. Cir. 1995), the court determined that the existence of a general method of identifying a specific DNA does not make the specific DNA obvious. Regarding structural or functional homologs, however, the court stated

"Normally, a *prima facie* case of obviousness is based upon structural similarity, i.e., an established structural relationship between a prior art compound and the claimed compound. Structural relationships may provide the requisite motivation or suggestion to modify known compounds to obtain new compounds. For example, a prior art compound may suggest its homologs because homologs often have similar properties and therefore chemists of ordinary skill would ordinarily contemplate making them to try to obtain compounds with improved properties (see page 9, paragraph 4 of attached ref)."

Since the claimed primers simply represent structural homologs, which are suggested by the prior art as useful for primers and probes, and concerning which a biochemist of ordinary skill would

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attempt to obtain alternate compounds with improved properties, the claimed primers and probes are *prima facie* obvious over the cited references in the absence of secondary considerations.

9. Claims 7 is rejected under 35 U.S.C. 103(a) as being unpatentable over Sargent et al (EMBO J. (1989) 8(8):2305-2312) in view of Beach et al (U.S. Patent 6,025,192).

Sargent teaches cosmid vectors which are transformed into E. Coli host cells (page 2311, column 2) which cosmid vectors comprise a double stranded nucleic acid (necessarily including sense and antisense strands) that includes the "G7" gene (page 2306, figure 1). Within figure 1, several cosmid vectors overlap the "G7" gene including F9N, F12M, FMC, FMEa and EL3. The "G7" gene is inherently found to be the MSH5 gene claimed. With regard to points of similarity, the "G7" gene is located on chromosome 6p21.3 as is the MSH5 gene, and above is listed the partial sequence comparison of MSH5 and "G7", where MSH5 is the Qy sequence and "G7" is the database sequence.

Sargent does not teach placement of the G7 gene into a retroviral vector.

Beach teaches placement of genes into retroviral vectors for the elucidation of mammalian gene function (abstract).

It would have been *prima facie* obvious to one having ordinary skill in the art at the time the invention was made to put the unknown G7 gene of Sargent into the retroviral vector of Beach since Beach states "The present invention relates to methods and compositions for the elucidation of mammalian gene function (abstract)". An ordinary practitioner would have been motivated to use the retroviral vector of Beach to identify the function of G7.

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Conclusion

10. Any inquiry concerning this communication or earlier communications from the examiner should be directed to Jeff Fredman, Ph.D. whose telephone number is (703) 308-6568.

If attempts to reach the examiner by telephone are unsuccessful, the examiner's supervisor, W. Gary Jones, can be reached on (703) 308-1152.

Any inquiry of a general nature or relating to the status of this application should be directed to the Technology Center receptionist whose telephone number is (703) 308-0196.

Papers related to this application may be submitted to Technology Center 1600 by facsimile transmission via the P.T.O. Fax Center located in Crystal Mall 1. The CM1 Fax Center numbers for Technology Center 1600 are either (703) 305-3014 or (703) 308-4242. Please note that the faxing of such papers must conform with the Notice to Comply published in the Official Gazette, 1096 OG 30 (November 15, 1989).



Jeffrey Fredman
Patent Examiner
Art Unit 1655

February 6, 2001